

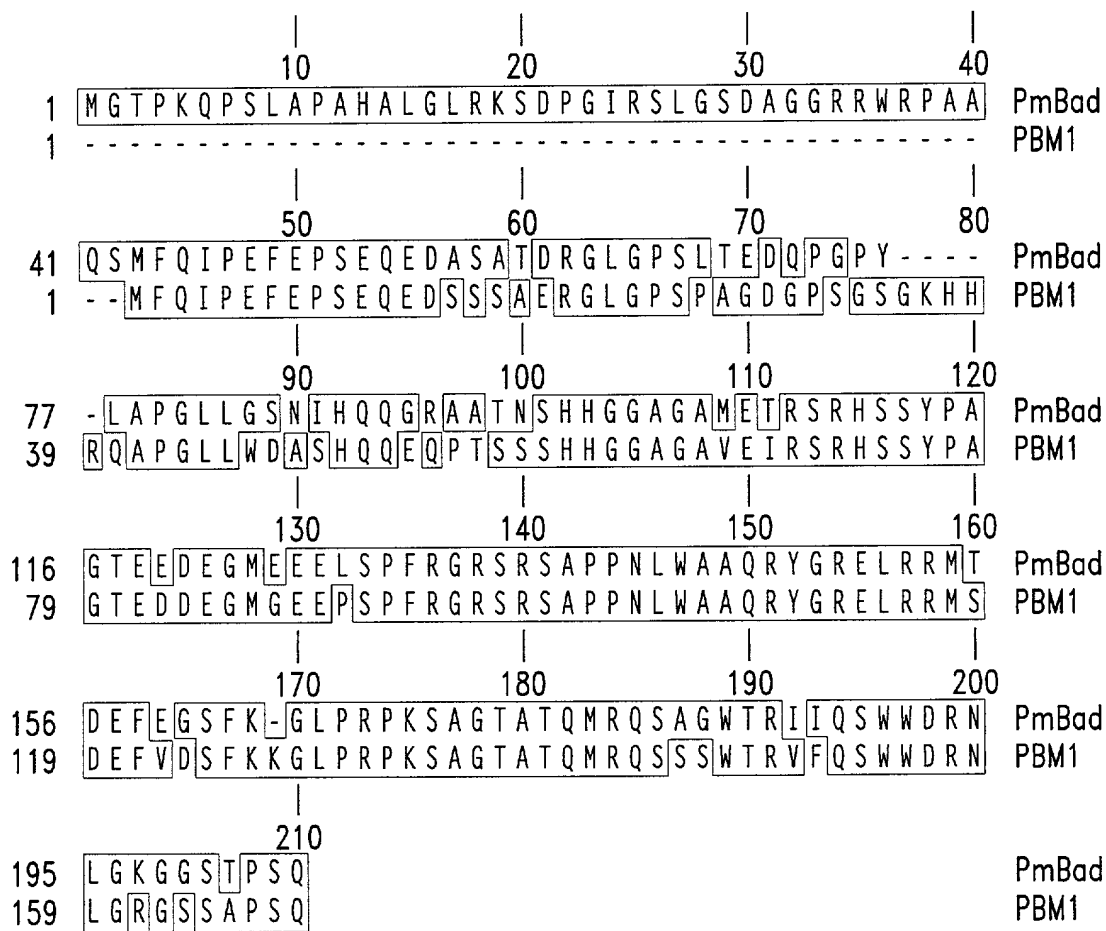
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Met Phe	
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Gln Ile Pro Glu Phe Glu Pro Ser Glu Gln Glu Asp Ser Ser Ser Ala	
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GAG AGG GGC CTG GGC CCC AGC CCC GCA GGG GAC GGG CCC TCA GGC TCC	153
Glu Arg Gly Leu Gly Pro Ser Pro Ala Gly Asp Gly Pro Ser Gly Ser	
20 25 30	
GGC AAG CAT CAT CGC CAG GCC CCA GGC CTC CTG TGG GAC GCC AGT CAC	201
Gly Lys His His Arg Gln Ala Pro Gly Leu Leu Trp Asp Ala Ser His	
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Gln Gln Glu Gln Pro Thr Ser Ser Ser His His Gly Gly Ala Gly Ala	
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Val Glu Ile Arg Ser Arg His Ser Ser Tyr Pro Ala Gly Thr Glu Asp	
70 75 80	
GAC GAA GGG ATG GGG GAG GAG CCC AGC CCC TTT CGG GGC CGC TCG CGC	345
Asp Glu Gly Met Gly Glu Glu Pro Ser Pro Phe Arg Gly Arg Ser Arg	
85 90 95	
TCG GCG CCC CCC AAC CTC TGG GCA GCA CAG CGC TAT GGC CGC GAG CTC	393
Ser Ala Pro Pro Asn Leu Trp Ala Ala Gln Arg Tyr Gly Arg Glu Leu	
100 105 110	
CGG AGG ATG AGT GAC GAG TTT GTG GAC TCC TTT AAG AAG GGA CTT CCT	441
Arg Arg Met Ser Asp Glu Phe Val Asp Ser Phe Lys Lys Gly Leu Pro	
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Arg Pro Lys Ser Ala Gly Thr Ala Thr Gln Met Arg Gln Ser Ser Ser	
135 140 145	
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Trp Thr Arg Val Phe Gln Ser Trp Trp Asp Arg Asn Leu Gly Arg Gly	
150 155 160	

*Fig. 1A*

*Fig. 1B*

[illegible]

# ALIGNMENT OF MOUSE AND HUMAN BAD PROTEIN SEQUENCES



PmBad = mouse protein sequence  
 PBM1 = human protein sequence

Fig. 2

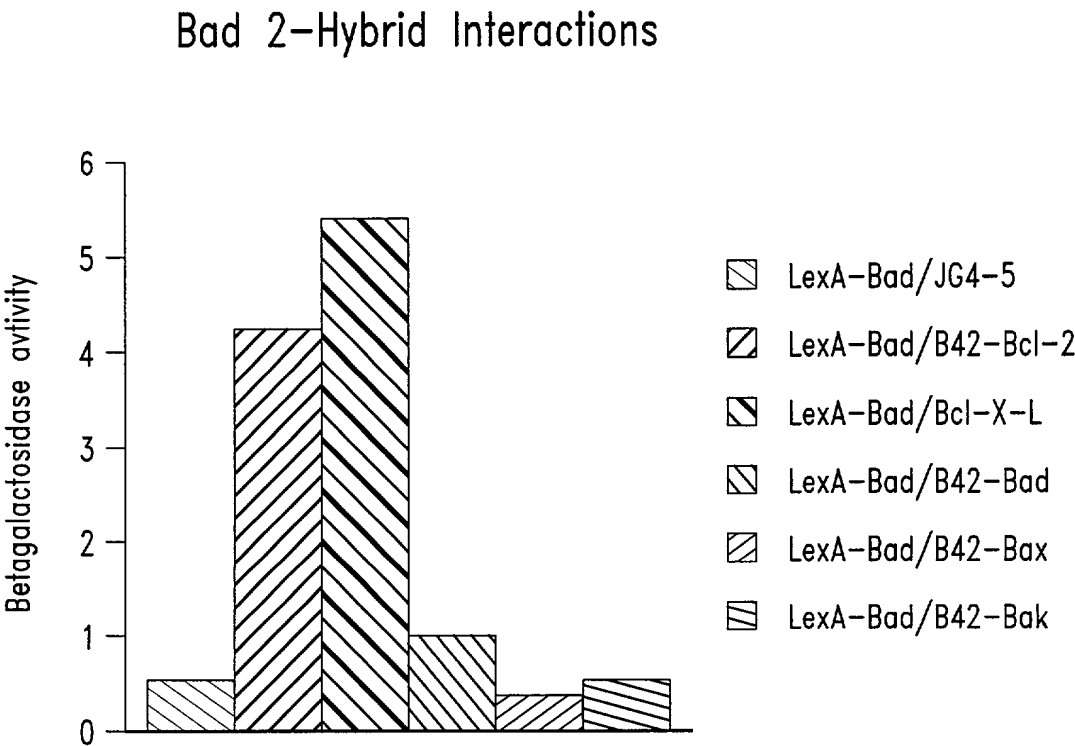


Fig. 3

Binding of Bcl-2 Family Proteins to Bad

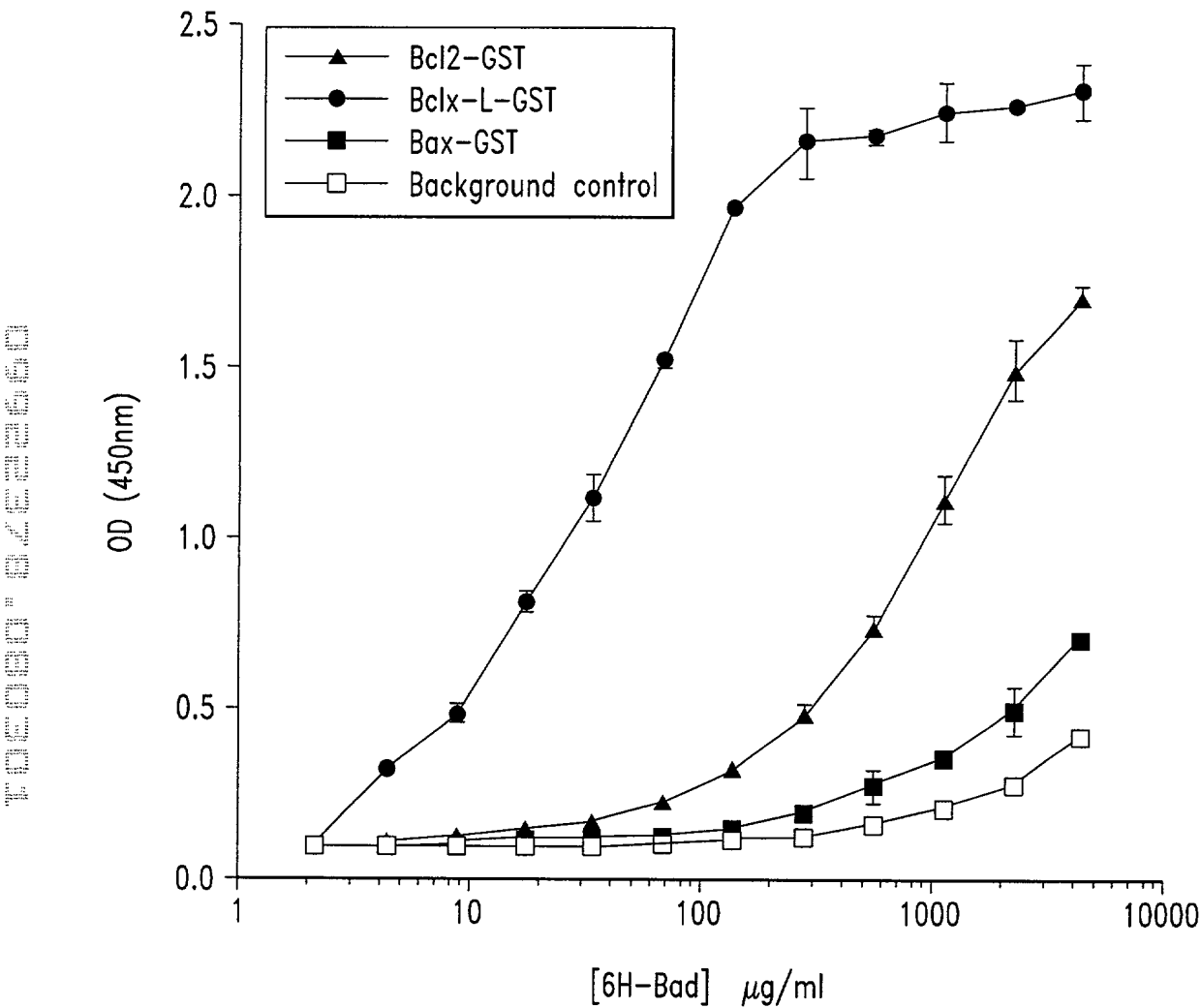


Fig. 4